Assistant Commissioner for Patents

Washington, D.C. 20231

Box: Non-Fee Amendment

Sir:

Transmitted herewith is an amendment (10 pages) and Appendix (5 pages) in the above-identified application. Also enclosed are:

- [] A Request for _____-Month Extension of Time.
- [] Applicant claims small entity status. (See 37 CFR 1.27.)
- [X] A self-addressed, prepaid postcard for acknowledging receipt.
- [X] No additional fee is required.
- [X] The fee has been calculated as shown below:

a) SMALL ENTITY

(Col. 1) (Col. 2) (Col. 3) b) LARGE ENTITY

	CLAIMS REMAINING AFTER AMENDMENT		HIGHEST NO. PREVIOUSLY PAID FOR	PRESENT EXTRA	RATE	ADD'L FEE
TOTAL	. 100	MINUS	100	0	a) \$ 9= b) \$18	\$0
INDEP	4	MINUS	4	0	a) \$40= b) \$80	\$0
· ·	SENTATION OF Time Fee (N	See fee Schedule	\$0			
					TOTAL	\$0

l J	A check for \$ is enclosed to cover the above fees.						
[]	Please charge my Deposit Account No. 14-1138 in the amount of \$ A duplicate copy of this sheet is enclosed.						
[X]	The Commissioner is hereby authorized to charge any additional fees or credit any overpayment to Deposit Account No. 14-1138. A duplicate copy of this sheet is enclosed.						
Date: Mar	Michael L. Goldman Registration No. 30,727						

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Certificate of Mailing - 37 CFR 1.8(a)

I hereby certify that this correspondence is being deposited with the United States Postal Service as first class mail in an envelope addressed to: Assistant Commissioner for Patents, Washington, D.C. 20231, on the date below.

Date /

Jane C. Wirszyla



Docket No.: 19603/468 (CRF D 1595C)

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants

Barany et al.

Serial No.

08/794,851

Filed

February 4, 1997

For

DETECTION OF NUCLEIC ACID

SEQUENCE DIFFERENCES USING THE

LIGASE DETECTION REACTION WITH

ADDRESSABLE ARRAYS

39/07

Art Unit:

1627

AMENDMENT UNDER 37 C.F.R. § 1.116

Assistant Commissioner for Patents

Washington, D.C. 20231

Box: Non-Fee Amendment

Dear Sir:

In response to the March 2, 2001, Advisory Action, please amend the above-identified patent application as follows:

In the Claims:

Please amend claims 1, 138, 150, and 151 as follows

(Four Times Amended) A method for identifying one or more of a plurality of sequences differing by one or more single-base changes, insertions, deletions, or translocations in a plurality of target nucleotide sequences comprising:

providing a sample potentially containing one or more target nucleotide sequences with a plurality of sequence differences;

providing a plurality of oligonucleotide probe sets, each set characterized by (a) a first oligonucleotide probe, having an oligonucleotide target-specific portion and an oligonucleotide addressable array-specific portion and (b) a second oligonucleotide probe, having an oligonucleotide target-specific portion and a detectable

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